## Euly. (3)

## 6. SEQUENCE LISTING

	(1) GENI	ERAL INFORMATION: APPLICANT: Gan, Z. R.			
5	*. (ii)		Chimeric Protein Containing An Intramolecular Chaperone-Like Sequence And Its Application To Insulin Production		
	(iii)	NUMBER OF SEQUENCES	. 7		
10	(iv)	CORRESPONDENCE ADDRESS:			
	(1*)	(A) ADDRESSEE:			
		(B) STREET:			
		(C) CITY:			
		(D) STATE:			
15		(E) COUNTRY:			
		(F) ZIP:			
	(v)	COMPUTER READABLE FORM:			
		(A) MEDIUM TYPE: 3.5 inch diskette			
		(B) COMPUTER: IBM	PC		
20		(C) OPERATING SYSTE			
		(D) SOFTWARE: Word			
	DATA:				
		•	MBER: To Be Assigned		
		(2)	Filed Concurrently Herewith		
25		(C) CLASSIFICATION;			
	(vii)	PRIOR APPLICATION DATA:			
		(A) APPLICATION NUM	ABER:		
		(B) FILING DATE:	The state of the s		
30	(viii)	ATTORNEY/AGENT INFO	RMATION:		
30		(A) NAME:	n (DED		
		(B) REGISTRATION NU			
		(C) REFERENCE/DOCK			
	(ix)	TELECOMMUNICATION 1	INFURMATION:		

TELEPHONE:

TELEFAX:

(A)

(B)

35

	1		
		(C) TELEX:	
1237	(2	) INFORMATION FOR SEQ ID NO: 1:	
/ الم كلو	•	(i) SEQUENCE CHARACTERISTICS:	
7		(A) LENGTH: 49 amino acids	
	5	(B) TYPE: amino acid	
		(C) TOPOLOGY: linear	
		(ii) MOLECULE TYPE: protein	
		(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
	M	Set Phe Pro Thr Ile Pro Leu Ser Arg Leu Phe Asp Asn Ala Met Leu	
	10 1	5 10	15
	A	rg Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe	
		20 25 30	
**	G	ilu Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Plie Leu Gln Asn	
		35 40 45	
	15 P	ro	
i. Sita		50	
70.		,	
- 74 - 122 - 151 - 151	(3	3) INFORMATION FOR SEQ ID NO: 2:	
		(i) SEQUENCE CHARACTERISTICS:	
ede for a	20	(A) LENGTH: 92 amino acids	
- 14 - 14		(B) TYPE: amino acid	
gr)		(C) TOPOLOGY: linear	
#1 - 13 - 25		(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
2.2	N	Met Phe Pro Thr Ile Pro Leu Ser Arg Leu Phe Asp Asn Ala Met Leu	
	25 1		15
	A	Arg Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe	
		20 25 30	
	C	Glu Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn	
		35 40 45	
	30 Ł	Pro Gln Thr Ser Leu Ser Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn	
		50 55 60	
	A	Arg Glu Glu Thr Gln Gln Lys Ser Asn Leu Glu Leu Leu Arg Ile Ser	

Leu Leu Ile Gln Ser Trp Leu Glu Pro Val Gln

John 1817

	(4) IN	IFORMATI	ON FOR S	EQ ID NO: 3	3:		
	(i)	SEQU	ENCE CHA	ARACTERIS	TICS:		
		(A)	LENGTH	: 6 amino a	cids		
		(B)	TYPE: a	mino acid			
5		(C)	TOPOLO	GY: linear			
	(x	i) SEQU	ENCE DES	SCRIPTION:	SEQ ID	NO: 3:	
	Leu Gly	Thr Gly Pro	) Arg				
	1		5				
LO (5) INFORMATION FOR SEQ ID NO: 4:							
	(i)	) SEQU	JENCE CH.	ARACTERIS	STICS:		
		(A)	LENGTH	: 86 amino	acids		
		<b>(B)</b>	TYPE: a	amino acid			
		(C)	TOPOLO	GY: linear			
15	()	(i) SEQUI	ENCE DES	CRIPTION:	SEQ ID	NO: 4:	
	Phe Val	Asn Gln Hi	s leu Cys G	ly Ser His Le	eu Val Glu	Ala Leu Tyr	
	1		5		10		15
	Leu Val	Cys Gly Gl	u Arg Gly F	Phe Phe Tyr	Thr Pro L	ys Thr Arg Arg	
		20		25			0
20	Glu Ala	Glu Asp Le	u Gln Val C	Gly Gln Val C	Glu Leu G	ly Gly Gly Pro	
		35		40		45	
	Gly Ala	Gly Ser Lei	ı Gln Pro L	eu Ala Leu C	du Gly Se	r Leu Gln Lys	
	5	0	55		60	)	
	Arg Gly	Ile Val Glu	Gln Cys C	ys Thr Ser Ile	e Cys Ser	Leu Tyr Gln	
25	65		70		75	80	
	Leu Glu	Asn Tyr C	ys <b>As</b> n				
			85	90			
	(6) I	NFORMAT	ION FOR S	SEQ ID NO:	5:		
30	(	i) SEQ	UENCE CH	IARACTERI	STICS:		
		(A)	LENGTH	I: 52 amino	acids		
		(B)	TYPE:	amino acid			
		(C)	TOPOLO	GY: linear			
	•	-		CRIPTION:			
35	Phe Val	Asn Gin H	is leu Cys C	Gly Ser His L	eu Val Gl	u Ala Leu Tyr	
	1		5		10		15
					29		

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Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr Arg Gly
                                   25
   Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu
                                                  45
                              40
 5 Asn Tyr Cys Asn
         50
                           55
         INFORMATION FOR SEQ ID NO: 6:
   (7)
                SEQUENCE CHARACTERISTICS:
         (i)
                      LENGTH: 107 amino acids
10
                (A)
                (B)
                      TYPE: amino acid
                      TOPOLOGY: linear
                (C)
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
   Met Phe Pro Thr Ile Pro Leu Ser Arg Leu Phe Asp Asn Ala Met Leu
                                                                        15
15 1
   Arg Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe
                                   25
                  20
   Glu Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn
                               40
20 Pro Leu Gly Thr Gly Pro Arg Phe Val Asn Gln His leu Cys Gly Ser
                            55
   His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe
                                                               80
                                           75
                         70
   65
   Tyr Thr Pro Lys Thr Arg Gly Ile Val Glu Gln Cys Cys Thr Ser Ile
                                                                    95
25
                     85
   Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
                                   105
                                                      110
                 100
          INFORMATION FOR SEQ ID NO: 7:
   (8)
30
                SEQUENCE CHARACTERISTICS:
          (i)
                (A)
                      LENGTH: 150 amino acids
                      TYPE: amino acid
                (B)
                      TOPOLOGY: linear
                (C)
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
35 Met Phe Pro Thr Ile Pro Leu Ser Arg Leu Phe Asp Asn Ala Met Leu
                                                                            15
                                                 10
                     5
    1
                                          30
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	Arg Ala His Arg Leu His Gin Leu Ala Phe Asp Thr Tyr Gin Giu Phe						
	20		25	30			
	Glu Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn						
	35	40		45			
5	Pro Gln Thr Ser Leu Ser Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn						
	50	55		60			
	Arg Glu Glu Thr Gln Gln Lys Ser Asn Leu Glu Leu Leu Arg Ile Ser						
	65	70		75		80	
	Leu Leu Ile Gln	Ser Trp Leu Gli	ı Pro Val Gl	n Leu Gly Thr Gly			
10	8:	5		90	95		
	Pro Arg Phe Val Asn Gln His leu Cys Gly Ser His Leu Val Glu Ala						
	100		105	110			
	Leu Tyr Leu Vai Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr						
	115		120		125		
15	5 Arg Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln						
	130	135		140			
	Leu Glu Asn Tyr Cys Asn						
	145	150					